



Fig.22

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1  atgaaataacctgctgccgaccgctgctgctggtctgctgctcctc
_M K Y L L P T A A A G L L L L
46  gctgcccagccggcgatggccatggacatcaagcataatcggccg
A A Q P A M A M D I K H N R P
91  cgcagagactgcgtggcagagggcaaagtgtgtgacctactgtgc
R R D C V A E G K V C D P L C
136 tcctctgggggatgctggggcccaggccctgggtcagtgtgtgtcc
S S G G C W G P G P G Q C L S
181 tgtcgaaattatagccgaggaggtgtctgtgtgacctactgcaac
C R N Y S R G G V C V T H C N
226 tttctgaatggggagcccctcgagcaccaccaccaccactga (SEQ ID NO:17)
F L N G E P L E H H H H H H (SEQ ID NO:14)

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Fig.23(SEQ ID NO:14)

ATGGTTTGTGTAGCCAGCTGTCCCCATAACTTTGTGGTGGATCAAACATCCTGTGTCA  
GGGCCTGTCCTCCTGACAAGATGGAAGTAGATAAAAATGGGCTCAAGATGTGTGAGC  
CTTGTGGGGGACTATGTCCCAAAGCCTGTGAGGGAACAGGCTCTGGGAGCCGCTTCCA  
GACTGTGGACTCGAGCAACATTGATGGATTTGTGAACTGCACCAAGATCCTGGGCAAC  
CTGGACTTTCTGATCACCGGCCTCAATGGAGACCCCTGGCACAAGATCCCTGCCCTGG  
ACCCAGAGAAGCTCAATGTCTTCCGGACAGTACGGGAGATCACAGGTTACCTGAACA  
TCCAGTCCTGGCCGCCCCACATGCACAACTTCAGTGTTTTTTTCCAATTTGACAACCATT  
GGAGGCAGAAAGCTTGCGGCCGCACTCGAGCACCACCACCACCACCTGA

Fig.24(SEQ ID NO :15)

Met V C V A S C P H N F V V D Q T S C V R A C P P D K Met E V D K N G L K Met C E P C  
G G L C P K A C E G T G S G S R F Q T V D S S N I D G F V N C T K I L G N L D F L I T G L  
N G D P W H K I P A L D P E K L N V F R T V R E I T G Y L N I Q S W P P H Met H N F S V  
F S N L T T I G G R ~~K L A A A L E H H H H H H~~  
S K L A A A L E H H H H H

Fig.25(SEQ ID NO :16)